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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/784,077

TIME: 14:45:34

DATE: 12/17/2001

Input Set : N:\Crf3\RULE60\09784077.txt
Output Set: N:\CRF3\12172001\1784077.raw

SEQUENCE LISTING

4 (1) GENERAL INFORMATION: (i) APPLICANT: NATSUKA, SHUNJI 6 7 GERSTEN, KEVIN M. 8 LOWE, JOHN B. (ii) TITLE OF INVENTION: MURINE ALPHA (1,3) FUCOSYLTRANSFERASE 10 11 FUC-TVII, DNA ENCODING THE SAME, METHOD FOR PREPARING THE 12 SAME, ANTIBODIES RECOGNIZING THE SAME, IMMUNOASSAYS FOR 13 DETECTING THE SAME, PLASMIDS CONTAINING SUCH DNA 15 (iii) NUMBER OF SEQUENCES: 4 17 (iv) CORRESPONDENCE ADDRESS: 18 (A) ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, 19 P.C. (B) STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400 20 ENTERED 21 (C) CITY: ARLINGTON 22 (D) STATE: VA 23 (E) COUNTRY: USA (F) ZIP: 22202 24 26 (V) COMPUTER READABLE FORM: 27 (A) MEDIUM TYPE: Floppy disk 28 (B) COMPUTER: IBM PC compatible 29 (C) OPERATING SYSTEM: PC-DOS/MS-DOS 30 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (vi) CURRENT APPLICATION DATA: 32 C--> 33 (A) APPLICATION NUMBER: US/09/784,077 (B) FILING DATE: 16-Feb-2001 C--> 3435 (C) CLASSIFICATION: 37 (vii) PRIOR APPLICATION DATA: 38 (A) APPLICATION NUMBER: US 08/613,098 39 (B) FILING DATE: 08-MAR-1996 44 (viii) ATTORNEY/AGENT INFORMATION: 45 (A) NAME: LAVALLEYE, JEAN-PAUL 46 (B) REGISTRATION NUMBER: 31,451 47 (C) REFERENCE/DOCKET NUMBER: 2363-114-55 49 (ix) TELECOMMUNICATION INFORMATION: 50 (A) TELEPHONE: 703-413-3000 (B) TELEFAX: 703-413-2220 54 (2) INFORMATION FOR SEQ ID NO: 1: (i) SEQUENCE CHARACTERISTICS: 56 (A) LENGTH: 3594 base pairs 57 58 (B) TYPE: nucleic acid 59 (C) STRANDEDNESS: double 60 (D) TOPOLOGY: linear (ii) MOLECULE TYPE: cDNA 62 67 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

69 ACAAACAGGA AGGACAGCAG GCTCTGGCAG CCAGAAGCCT GTGGCCCCAA GCTGGCAGGA

71 TGGCCCCCTT CCTGCAGGTC CCCCACAGCC TTCTGGGTTC CTGACACGAG AGAAGAGGTG

60

120

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| 73 | GGGCGGGGTG | AAGTGAACTC | TGAAGCCAAA | ATGTGACTCT | CCTGGGGTCA | CCAGCTTGGG | 180 |
|-----|------------|--------------|------------|-------------|--------------|------------|------|
| 75 | GAGAGGTGAA | GAAAGATGCC | GGGGCGGAAA | CAAAGGGGCA | GATATCACTA | TGGTTATCTT | 240 |
| 77 | ACTAAGCACA | GAGTAACTGA | AAAAGCAAGG | GTACCGCTGC | CCACCTCGTG | CCCACCTTAC | 300 |
| 79 | GTTATACCTC | AAACCAGCTA | GATAGTTTCT | GATGGCACCC | ATACCCTCCC | TTCCCCTTTA | 360 |
| 81 | GGCATTGCGC | AAGCTCTCCA | CCACAATCTG | GAAGTTATAC. | CCTGCGAGGG | GATGGGCAGG | 420 |
| 83 | GCACTTCTGA | GGTGCCAATC | AGCCTGCACT | CGCCTCTGCC | CTGGCCATGG | CACTGCTGTC | 480 |
| 85 | AGTTTCTTGG | TACCTGTCTC | AACAGCAGCC | TTGTCACGTG | AGACTATGGC | TGGCGGTGGG | 540 |
| 87 | GGTGGGGGCA | GGAATCCTAG | AAGCACAGGA | GTGACATAGG | GTCGGGTCGG | GCAGAGCGAA | 600 |
| 89 | GTGTAGGAGG | TGATCCCCAA | AGGGATGCTG | GGGACGATCT | GGCCAACACT | GTCCTCCCAT | 660 |
| 91 | TCAAAACTCC | CAGTCTGGAG | CTCTGGGACA | TGGACAAGCC | AGGCCTGCTA | TTCTCCATAC | 720 |
| 93 | AGGGCTCCAT | AGTGTCTGGC | TCAGCAGAGT | GGGGGATCTG | GTGGGGATGG | AGGAAGCTTA | 780 |
| 95 | GCTAAAAGCT | TTGTATAGGC | TGAAGCTCTG | AGTGACCCTG | CTGGGCCACC | CTACCCTGGT | 840 |
| 97 | CTGGGCTGGG | TCATTGCATC | CCCAGATTGG | AAGGCTTGGT | GAGATGGAGA | GGAACCTTGG | 900 |
| 99 | CTACAAGCTA | TAGCTTTGCC | CACCAGAGCC | TGCTGGAGGG | GAATCAAACA | AGCCTGGACC | 960 |
| 101 | TGAGGCTGG | ACTAGCTTTC | CTGTTTCTGG | AGTGGATGC | AACCCCCTGC | CCACCAGCCT | 1020 |
| 103 | GCCTGTCCAC | CGCCAGGGACA | CACAGACTCC | TTCCCTTTCC | CAGACTGGAAA | GCCCCCTCCT | 1080 |
| 105 | GGGAGAGCAG | GAAGGAAGCA | ACCTGCAACT | CTTCCAGCCC | TGGACCTTGG | GCTGAACCTA | 1140 |
| 107 | CAGTTCAAGG | TTTGTATGCT | CACAGGTCTT | GGCAGGGAA | A GATAAGAATC | CCCAGGGCAC | 1200 |
| 109 | CCTCCCCCC | GCCCCCAGT | CCACTGCAGG | TAGCTCCTGC | GTCTGCCCTT | CAGGGCAAGT | 1260 |
| 111 | GCTGACGCTC | CATCAGACTG | TGATGGGGCC | CTTTTCTGAG | GATGACAATT | CTGAGAACAA | 1320 |
| 113 | GGCATTTTTC | TAGAGGTGGC | AGAACAGCAT | TTTGTGATGC | CCGAGGATCT | GGGAGCACAG | 1380 |
| 115 | GTCCAGCTTA | ATGAGGGATT | GGAGGAAGTG | GGTATCATCA | A TTACAGGGAG | GGGCCTCTGT | 1440 |
| 117 | GGCCTCCTGG | GAAAATGCAG | TTGCTCTCTT | TGGGTGGCCT | GGGGTTGTGT | GGTGGGCAGA | 1500 |
| 119 | GGACGGAGGT | GCTCATTGGG | GGAAGGGATC | ACTTCTGCTC | CAGAGTGCTCG | CAAGGGCCTT | 1560 |
| 121 | TCCTTTTCCT | GAAGGCAAGC | AGGCCTCCTC | CTCCTCCTCT | TCCTCCTTCT | CCTCTTCCTC | 1620 |
| 123 | CTCTTTCTCC | ATATGCCTAG | CTGGTCATTT | CTAGGGACCA | GCATGGTTGG | GAAGGGGCC | 1680 |
| 125 | TTGTCTTGGC | CTTCCTCTTG | TCTCAATTCC | CTCTTTGAGO | AGAAGACGGG | GTGGGTGGGG | 1740 |
| 127 | TAGGATTGGA | A TAGTGGTTGA | TGCCAAAGAT | TGAAGGGGTA | GGGCGGGGCA | GAAGTGGGAA | 1800 |
| 129 | GGTCCCTGGC | TTCCTCACCT | TGGTAGATGG | TGAGGAGCCC | CAGAGGTTGA | GCTGAGCAGC | 1860 |
| 131 | AGCTGTGATT | TCAGGGTGCC | TCTGTTGGAG | AGGCTGCTGT | GATTTGAAAA | TCTTCTTTCC | 1920 |
| 133 | TTGGTGACAA | TTCCAGAAGG | CTCCAGATGA | ATTGTATTGG | TGAGTGCCTG | GCCCTTAAGC | 1980 |
| 135 | AGTCCCAGCT | GGGGATGATG | GGGATTTATG | GGTGTCCCTG | AGCCTAGGGT | GACAGGGCCT | 2040 |
| 137 | CTCCTTTTT | TTTTATTCTG | CTTCAGGGTA | CCACCCCACC | AGGAGGCTGC | GGGCCTGGGG | 2100 |
| | | | | | TTCTGGCTGT | | 2160 |
| | | | | | CTTATCTGGC | | 2220 |
| 143 | CACCAACCGG | CCGCCAGAGC | TACCTGGTGA | CACCTGCACT | CGCTATGGCA | TGGCCAGCTG | 2280 |
| | | | | | GCTGTGGTCT | | 2340 |
| | | | | | AGGCCACACG | | 2400 |
| | | | | | CTCCATCGCT | | 2460 |
| | | | | | TTTGTACCCT | | 2520 |
| | | | | | AGCAGGATGG | | 2580 |
| | | | | | TACCGGCAGC | | 2640 |
| | | | | | CTATGCGCTA | | 2700 |
| | | | | | AACTCACAGC | | 2760 |
| | | | | | GCTGTACCCG | | 2820 |
| | | | | | GCCTTTGTAC | | 2880 |
| | | | | | ATGAATGAGA | | 2940 |
| | | | | | CTGGGTGACT | | 3000 |
| 169 | CTTCTGCACC | : ATCTGTGCCC | GCTACCCTTA | CTTGCCCCGC | AGCCAGGTCT | ATGAAGACCT | 3060 |

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| 173 175 177 179 181 183 185 | TGAAAGCTGG TTCCAGGCTT GAACTCCTGC TGCTGGGAGA GGCTGGATGG GTGGGAGACT GATGTTGAAA CCAAAGAGCT GGGCATCCAG GCTTTTGGTC ACCATGGCAC TACCCCAAGG CTTTTCCTGT TCAGTGAGCA GGAATTCAGG ATATAAGGAG AAGACTGGGC TGAGATACCC TGGTGGGCTT TAGAGTAGGG GCCCAGGATA AGAGACAATG AATTAATGAG GAGCATATGG GGAAGGTGGC TGAGGTCCC TGACTTACCT TGACCCATGG CTGAAGGCTC CATGCCCATG GCTGGAGCTG GGACCCTACA CTTCTATAGT CAAGGTGCTT AGCCTCAAGG TTGCAGATGC ACCCTCTAGT ACTCTGGGTG CAGACTGTAC ACTGGGCGCA GGGGGTTGTG GAAGGACAGT GCAGATGATT CTGGGCTTTT GACACCACAG TTCCCCCAGG GAAAGAGGCA CTACTAATAA AAACACTGAC AGAAATCTCC TGGTCAAGTC TGTTAGGCAG CAGAGCTCGA ATTC (2) INFORMATION FOR SEQ ID NO: 2: | | | | | | | | | | | | | G G G C | 3120 3180 3240 3300 3360 3420 3480 3540 3594 | | |
|---|--|------------|--------|------------|------------|--------|----------|----------------|----------|------|------------|-----------|-----------|------------------|--|------|--|
| 191 | (i) | SEQ | UENC: | E CH | ARAC | TERI | STIC | S: | | | | | | | | | |
| 192 | · · | | | | | | | | | | | | | | | | |
| 193 | | | | | | | | | | | | | | | | | |
| 194 | · , , | | | | | | | | | | | | | | | | |
| 195 | (D) TOPOLOGY: linear | | | | | | | | | | | | | | | | |
| 197 | (ii) MOLECULE TYPE: protein | | | | | | | | | | | | | | | | |
| 202 | (xi) | | | | | | | | | | | | | | | | |
| 204 | | Pro | Thr | Pro | _ | Pro | Pro | Ala | Cys | | Ser | Thr | Pro | Gly | | His | |
| 205 | 1 | | | | 5 | | | | | 10 | _ | _ | _ | | 15 | _ | |
| 207 | Arg | Leu | Leu | | Phe | Pro | Asp | \mathtt{Trp} | | Ala | Pro | Ser | Trp | | Ser | Arg | |
| 208 | . | a 1 | | 20 | a | | a | a | 25 | D | a 1 | D | | 30 | a 1 | D | |
| 210 | ьys | GIU | | | Cys | Asn | ser | ser 40 | ser | Pro | GIY | Pro | | Ата | GIU | Pro | |
| 211 213 | Пhr | Wa l | 35 · | | Acn | Cvc | Tlo | | m | Uic | Dro | Thr | 45 | λνα | Lou | λκα | |
| 213 | 1111 | 50 | GIII | Met | ASII | Cys | 55 | СТУ | TAT | urs | PIO | Thr 60 | AIG | ALG | Leu | Alg | |
| 214 | Δla | - | Glv | Glv | T.011 | Δla | | Glv | Δla | Thr | Dhe | Met | Va l | T۱۵ | Trn | Phe | |
| 217 | 65 | 111 | 011 | 017 | Dea | 70 | 011 | 0-1 | 71.Lu | | 75 | 1100 | , , , | | | 80 | |
| 219 | | Trp | Leu | Trp | Glv | | Ala | Pro | Glv | Ser | | Pro | Val | Pro | Gln | | |
| 220 | | | | | 85 | | | | | 90 | | | | | 95 | | |
| 222 | Thr | Leu | Thr | Ile | Leu | Ile | Trp | His | Trp | Pro | Phe | Thr | Asn | Arg | Pro | Pro | |
| 223 | | | | 100 | | | _ | | 105 | | | | | 110 | | | |
| 225 | Glu | Leu | Pro | Gly | Asp | Thr | Cys | Thr | Arg | Tyr | Gly | Met | Ala | Ser | Cys | Arg | |
| 226 | | | 115 | | | | | 120 | | | | | 125 | | | | |
| 228 | Leu | | Ala | Asn | Arg | Ser | | Leu | Ala | Ser | Ala | Asp | Ala | Val | Val | Phe | |
| 229 | | 130 | | _ | | _ | 135 | | _ | | | 140 | | _ | | | |
| 231 | | His | Arg | Glu | Leu | | Thr | Arg | Gln | Ser | | Leu | Pro | Leu | Asp | | |
| 232 | 145 | n | ** 1 _ | a 1 | ~ 1 | 150 | | ** - 1 | ~ | | 155 | 34-4 | 01 | | D | 160 | |
| 234 | Arg | Pro | HIS | GTÄ | | Pro | Trp | vaı | Trp | | ser | Met | GIU | ser | | ser | |
| 235 237 | λan | mh r | uic | C1., | 165 | II i o | 7 ×~ | Dho | A === | 170 | T10 | Phe | λan | Trn | 175 | T OU | |
| 238 | ASII | 1111 | птэ | 180 | Leu | птэ | Ary | Pile | 185 | GTÄ | ire | Pile | ASII | 190 | vai | Leu | |
| 240 | Sor | Тτε | Δτα | | Δen | Sar | λen | т1Д | | Va 1 | Dro | Tyr | Glv | | T.e.u | Glu | |
| 241 | 261 | - 1 - | 195 | nr 9 | пэр | JEI | asp | 200 | : 11C | *41 | 110 | - Y - | 205 | **** 9 | Leu | Jau | |
| 243 | Pro | Leu | | G] v | Pro | Thr | Ser | | Leu | Pro | Ala | Lys | | Ara | Met | Ala | |
| 244 | 110 | 210 | | 1 | 0 | | 215 | | | | | 220 | | 3 | | | |
| 246 | Ala | | Val | Ile | Ser | Asn | | Gln | Glu | Arg | Gln | Gln | Arg | Ala | Lys | Leu | |
| 247 | 225 | - | | | | 230 | | | | , | 235 | | _ | | - | 240 | |
| 249 | | Arg | Gln | Leu | Ala | Pro | His | Leu | Gln | Val | Asp | Val | Phe | Gly | Arg | Ala | |
| 250 | | | | | 245 | | | | | 250 | | | | | 255 | | |
| | | | | | | | | | | | | | | | | | |

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| 252 | Ser | Gly | Arg | | Leu | Cys | Ala | Asn | _ | Leu | Leu | Pro | Thr | Leu | Ala | Arg | |
|------------|---------------------------------------|-----|-----|-----|----------------|-----|-----|---------------|----------------|-------|-------|-------|-----|-----|----------|-----|----|
| 253 | | | _ | 260 | | _ | | _ | 265 | | _ | | | 270 | _ | | |
| 255 | Tyr | Arg | | Tyr | Leu | Ala | Phe | | Asn | Ser | Gln | His | _ | Asp | Tyr | Ile | |
| 256 | _ | | 275 | | • | | | 280 | _ | | | | 285 | | | | |
| 258 | Thr | | Lys | Phe | \mathtt{Trp} | Arg | | Ala | Leu | Ala | Ala | | Ala | Val | Pro | Val | |
| 259 | _ | 290 | _ | | | | 295 | _ | | | | 300 | | _ | _ | _ | |
| 261 | | Leu | Gly | Pro | Pro | _ | Ala | Thr | Tyr | Glu | | Phe | Val | Pro | Pro | _ | |
| 262 | 305 | _ | _ | | _ | 310 | | | _ | _ | 315 | _ | | _ | | 320 | |
| 264 | Ala | Phe | Val | His | | Asp | Asp | Phe | Ser | | Ala | Arg | Glu | Leu | | Val | |
| 265 | _ | | _ | | 325 | | _ | | | 330 | | | | | 335 | | |
| 267 | Phe | Leu | Val | | Met | Asn | Glu | Ser | | Tyr | Arg | GLY | Phe | Phe | Ala | Trp | |
| 268 | | | | 340 | | | | _ | 345 | | | _ | _ | 350 | _ | _, | |
| 270 | Arg | Asp | _ | Leu | Arg | Val | Arg | | Leu | Gly | Asp | Trp | _ | Glu | Arg | Phe | |
| 271 | | | 355 | | _ | | | 360 | _ | | | | 365 | | - | _ | |
| 273 | Cys | | Ile | Cys | Ala | Arg | _ | Pro | \mathtt{Tyr} | Leu | Pro | | Ser | Gln | Val | Tyr | |
| 274 | | 370 | | | | _ | 375 | | | | | 380 | | ٠ | | | |
| 276 | | Asp | Leu | Glu | Ser | - | Phe | Gln | Ala | | | | | | | | |
| 277 | 385 | | | | | 390 | | | | | | | | | | | |
| | 9 (2) INFORMATION FOR SEQ ID NO: 3: | | | | | | | | | | | | | | | | |
| 281 | , , , , , , , , , , , , , , , , , , , | | | | | | | | | | | | | | | | |
| 282 | • | | | | | | | | | | | | | | | | |
| 283 | | | | | | | | | | | | | | | | | |
| 284 | | | | | | | _ | Le | | | | | | | | | |
| 285 287 | (44) | • | | | GY:] | | | منداد | | | | | | | | | |
| 288 | (ii) | | | | PE: C | | | | | | יםם י | - MED | • | | | | |
| 293 | (xi) | | | | | | | | | | . PK. | LMEK | | | | | |
| | GCGCGGAT | | | | | | | | | | - | | | | | | 41 |
| | (2) INFO | | | | | | | | -111(| JAC (| - | | | | | | 47 |
| 299 | | | | | ARACI | | | | | | | | | | | | |
| 300 | (1) | | | | : 44 | | | | | | | | | | | | |
| 301 | | • • | • | | nucle | | - | 113 | | | | | | | | | |
| 302 | | • • | , | | EDNES | | | ۵ ا | | | | | | | | | |
| 303 | | | | | SY: 1 | | _ | | | | | | | | | | |
| 305 | (ii) | | | | | | | elei <i>c</i> | r aci | Ы | | | | | | | |
| 306 | () | | | | | | | | | | PRI | MER' | , | | | | |
| 311 | · · | | | | | | | | | | | | | | | | |
| | GCGCGGAT | | | | | | | | | | TTC | | | | | | 44 |
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VERIFICATION SUMMARY

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Output Set: N:\CRF3\12172001\1784077.raw

L:33 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:34 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]